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Sequence 39592, A Sequence 31796, A Sequence 31096, A Sequence 4034, Ap Sequence 1120, Ap Sequence 1120, Ap Sequence 123, App Sequence 223, Appl Sequence 2239, Ap Sequence 2243, Ap Sequence 2017, Ap Sequence 2019, Ap Sequence 2019, Ap Sequence 2189, Ap Sequence 22890, A Sequence 22890, A Sequence 22890, Ap Sequence 119, Appl Sequence 12890, A Sequence 22890, A Sequence 48, Appl
                                                                                                                                     February 16, 2003, 17:00:49 ; Search time 193.217 Seconds (without alignments) 10829.323 Million cell updates/sec
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Copyright (c) 1993 - 2003 Compugen Ltd.
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US-10-203-138A-8641
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US-10-203-138A-3500
US-10-144-771-4034
US-10-02-411A-1120
US-10-09-411A-1120
US-10-017-161-1913
US-10-104-771-4707
US-10-104-771-4707
US-10-104-771-4707
US-10-104-771-229
US-10-109-411A-2243
US-10-104-771-2049
US-10-11-161-2047
US-10-17-161-2049
US-10-203-138A-8573
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US-10-144-771-2890
US-10-203-138A-8573
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US-10-203-138A-227
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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22103, Ap
2245, Ap
21113, Ap
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                                                                                                                                                                                                                                                                                                                                                                                                                     282 ATGCCCTGCTGGTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGA 341
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                462 CGCTACCATAGTCGCATAATGTAACGTCGCATGTCCTACTGGTACTGCACTTGATGATGG 521
            Sequence 5
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                                                                                                                                                                                                                                       Sequence 39592, Application US/10144771
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig
TILLE OF INVENTION: HUMAN GENOME DISCOVERY SYSTEM AND USES THEREOF
CURRENT FILLNG DATE: 2002-05-15
UNRENT FILLNG DATE: 2002-05-15
NUMBER OF SEQ ID NOS: 47235
SEQ ID NO 39592
LENGTH: 897
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         US-60-434-832-6682

US-10-144-771-33054

US-09-806-708B-22

US-09-531-113-35330

US-09-950-084-2103

US-10-017-161-2245

US-10-144-771-21113

US-10-017-161-2245

US-09-950-084-771-21113

US-09-950-084-771-21113

US-09-950-084-7446

US-09-950-084-7446

US-09-950-084-7446

US-09-950-084-7446

US-09-724-676-30403

US-09-724-676-30403

US-09-724-676-30402
                                                                                                                                                                                                                                                                                                                                                                      5.5%; Score 77.4; DB 6; 43.9%; Pred. No. 3.4e-08; ative 0; Mismatches 421;
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                                                                                                                                                                                                                                                                                                                                                                                       Matches 330; Conservative
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US-10-144-771-39592
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TYPE: DNA
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                                           702 CGCATAATGTAACGTTGCATGCCCTGATGGTACTATAAGTGCTGCTGGAGTAAATAG
                                                                                                        762 GGTAGCACAAAACACTGAATGTACTAATTGTGCTCCTAACTTTTACAATAATAATGCTCC
                                                                                                                          TAATTTCAATCCAGGTAATAGTACATGCCTACCTTGCCCAGCAAATAAAGATTATGGTGC
Sequence 31796, Application US/10144771
Sequence 31796, Application US/10144771
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig
TITLE OF INVENTION: HUMAN GENOME DISCOVERY SYSTEM AND USES THEREOF
FILE REFERENCE: CL00131
CURRENT APPLICATION NUMBER: US/10/144,771
CURRENT FILING DATE: 2002-05-15
NUMBER OF SEQ ID NOS: 47235
SEQ ID NO 31796
LENGTH: 961
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Best Local Similarity 43.2
Matches 312; Conservative
                                                                                                                                                                                                                                                                                                                                                                                RESULT 2
US-10-144-771-31796/c
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; ORGANISM: HUMAN
US-10-144-771-31796
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APPLICANT: Chen, Wensheng
TITLE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
FILE REFERENCE: PB 0004 WO 8
CURRENT APPLICATION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO FILE REFERENCE: PB 0004 WO 8
CURRENT APPLICATION NUMBER: US/10/203,138A
CURRENT APPLICATION NUMBER: US 60/203,312
PRIOR PILING DATE: 2002-08-02
PRIOR FILING DATE: 2004-06
PRIOR FILING DATE: 2004-06
PRIOR FILING DATE: 2004-06
PRIOR FILING DATE: 2004-06
PRIOR PRIOR APPLICATION NUMBER: US 60/203,366
PRIOR FILING DATE: 13 Aduatt 2000 (33.08.00)
PRIOR PRIOR APPLICATION NUMBER: US 60/203,466
PRIOR PILING DATE: 21 September 2000 (31.08.00)
PRIOR PRILING DATE: 21 September 2000 (21.09.00)
PRIOR PILING DATE: 21 September 2000 (21.09.00)
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PILING DATE: 21 September 2000 (21.09.00)
PRIOR APPLICATION NUMBER: US 60/234,687
PRIOR PRIOR PLING DATE: 21 DATE
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Length 1635;
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ilarity 41.5%; Pred. No. 0.0003;
Conservative 0; Mismatches 650; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Molecular Dynamics Sequence Listing Engine SEQ ID NO 8641 LENGTH: 1635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = US-10-203-138A-8641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-10-203-138A-8641/c; Sequence 8641, Application US/10203138A; GENERAL INFORMATION:
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APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Penn, Sharron G.
Rank, David R.
Hanzel, David K.
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APPLICANT: FenD, SHALTON G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: Hanzel, David K.
APPLICANT: HANZEL, DAVID K.
APPLICANT: HUMBLE G.
TITLE OF INVENTION: HUMBN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FC
TILLE REFERENCE: PB 0004 MO 8
CURRENT APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/180,312
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 20 May 2000 (26.05)
PRIOR FILING DATE: 20 May 2000 (26.05)
PRIOR FILING DATE: 03 Adust 2000 (03.10.00)
PRIOR FILING DATE: US 60/233,366
PRIOR FILING DATE: US 60/233,467
PRIOR FILING DATE: US 60/234,487
PRIOR FILING DATE: 27 September 2000 (27.09.00)
PRIOR FILING DATE: 21 September 2000 (27.09.00)
PRIOR FILING DATE: 30 JUME 2000 (30.06.00)
PRIOR FILING DATE: 31 September 2000 (31.09.00)
PRIOR FILING DATE: 30 JUME 2000 (30.06.00)
SOUTH APPLICATION NUMBER: US 90/608,408
SOUTH APPLICATION NUMBER: US 
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4.3%; Score 60; DB 6; Length 1973;
Best Local Similarity 41.5%; Pred. No. 0.00031;
Matches 463; Conservative 0; Mismatches 650; Indels
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Sequence 3500, Application US/10203138A GENERAL INFORMATION:
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                                                              APPLICANT: Molecular Dynamics, Inc
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ORGANISM: Homo sapiens
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TITLE OF INVENTION: HUMAN GENOME DISCOVERY SYSTEM AND USES THEREOF
FILE REFERENCE: CLOOJ321
CURRENT APPLICATION NUMBER: US/10/144,771
CURRENT FILING DATE: 2002-05-15
NUMBER OF SEQ ID NOS: 47235
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Pred. No. 0.0088;
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: Sequence 4034, Application US/10144771
; GENERAL INFORMATION:
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ORGANISM: HUMAN
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APPLICANT: Chen, Wensheng
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXPRESSION IN BT 474
TITLE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
TITLE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
TITLE OF INVENTION: ANALYSIS OF GENE EXPRESSION IN BT 474
CURRENT APPLICATION NUMBER: US/10/203,138A
CURRENT FILING DATE: 2002-08-02
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PELING DATE: 04 February 2000 (04.02.00)
PRIOR PELING DATE: 03 August 2000 (25.05.00)
PRIOR PELING DATE: 03 August 2000 (03.08.00)
PRIOR PELING DATE: 03 August 2000 (03.08.00)
PRIOR PELING DATE: 1 September 2000 (21.09.00)
PRIOR PELING DATE: 2 September 2000 (21.09.00)
PRIOR PELING DATE: 2 September 2000 (21.09.00)
PRIOR PELING DATE: 2 September 2000 (30.08.00)
PRIOR PELING DATE: 2 September 2000 (30.08.00)
PRIOR PELING DATE: 3 September 2000 (30.08.00)
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                                                                                                                                                                                                        CCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACAGAATGTGTTAATTGTA 355
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                                                                                CAAATCCACCTGCTACTGCTAATTTAGTCACATAATGTAACGTTAAATGCCCTGCTGGTA
Sequence 7146, Application US/10203138A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Molecular Dynamics, Inc.
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Rank, David R.
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US-10-203-138A-7146/c
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APPLICANT:
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LOCATION: (1)..(1141)
COTHER INFORMATION: Consensus sequence of A.t., L.a., and B.n. FAEI promoters US-09-806-708B-22
                    487 GTCGCATGTCCTACTGGTACTGCACTTGATGGAGTAACTACTGATTATGTTAGATCA 546
                                                                                                                                                                                                                                                                                                                                 193 TCTGGCGATGCACATCATATACAGCACCTGCGCTGAAGTGAAGGTGGCTCACGAGCT 852
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GENERAL INFORMATION:
APPLICANT: The University of British Columbia
TITLE OF INVENTION: Requiation of Embryonic Transcription in Plants
FILE REFERENCE: 4810-56741
CURRENT APPLICATION NUMBER: US/09/806,708B
CURRENT FILING DATE: 2001-04-03
PRIOR APPLICATION NUMBER: US 60/147,133
PRIOR FILING DATE: 1999-08-04
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SOFTWARE: PatentIn version 3.0
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ORGANISM: Artificial sequence
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US-09-806-708B-22
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
TITLE OF INVENTION: BEIDERHIDIS FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/10/092,411A
PRIOR PELING DATE: 1098-08-13
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1998-08-13
PRIOR FILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-11-08
PRIOR FILING DATE: 1997-08-14
SEQ ID NOS: 5676
LENGTH: 1248
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3.6%; Score 50; DB 6; Length 1248;
Best Local Similarity 42.8%; Pred. No. 0.052;
Matches 254; Conservative 0; Mismatches 340; Indels
                                                                                                                                                                                                                              Query Match 3.7%; Score 52.2; DB 6; Length 5 Best Local Similarity 46.2%; Pred. No. 0.014; Matches 174; Conservative 0; Mismatches 203; Indels
                                                                                                                                                                       ; OTHER INFORMATION: NT HIT: U01287.1, EVALUE 1.30e+00 US-10-203-138A-7146
                                                                                       OTHER INFORMATION: MAP TO AL035457.11
FEATURE:
OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL = 4.6
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US-10-092-411A-1120
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                                                       ORGANISM: Homo sapiens
FEATURE:
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SEQ ID NO 7146
LENGTH: 510
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                                                                                                                  GGTAAACAGAGTIGGIGGIGGIGCATIGACIGCIGGIAAIGCCGCIACCAIAGICGCAIAAIG 482
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139 HHYVTAMNNAWTTMCMMDKDDKRTRWWWKKNNNATGWDDDTKYHMWNNGCBTVTWMVRY
                                                        KTDRDWSBKRMNYGMBWWKNWSYDVTYYWWVWDDMCKRKVRRWYRTRGRMRNYMVAWBTA
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Sequence 1913, Application US/10017161
GENERAL INFORMATION:
APPLICANT: SUWA, MAKIKO
APPLICANT: ASAI, KIYOSHI
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US-10-144-771-4707/c
Sequence 4707, Application US/10144771
GENERAL INFORMATION:
APPLICATIVE VENTER, J. Craig
TITLE OF INVENTION: HUMAN GENOME DISCOVERY SYSTEM AND USES THEREOF
FILE REFERENCE: CL001321
CURRENT APPLICATION NUMBER: US/10/144,771
CURRENT FILING DATE: 2002-05-15
UNMBER OF SEQ ID NOS: 47235
SEQ ID NO 4707
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               APPLICANT: ABURTANI, HIROYUKI
TITLE OF INVENTION: NOVEL G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 084335/0152
CURRENT APPLICATION NUMBER: US/10/017,161
CURRENT FILING DATE: 2002-12-18
PRIOR APPLICATION NUMBER: JP 2001/246789
PRIOR FILING DATE: 2001-06-18
NUMBER OF SEQ ID NOS: 2430
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. 0.17;
0; Mismatches 260;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 48;
Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3.4%;
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Best Local Similarity 44.0%
Matches 204; Conservative
AKIYAMA, YUTAKA
                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; NAME/KEY: CDS
; LOCATION: (2622)..(3332)
US-10-017-161-1913
                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: CDS
LOCATION: (201)..(437)
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LENGTH: 3532
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                                                                                                                                                                                                                                                             TYPE: DNA
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APPLICANT: MOJECULAR Dynamics, Inc.
APPLICANT: MOJECULAR Dynamics, Inc.
APPLICANT: MOJECULAR DAVID G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
APPLICANT: Chen, Wensheng
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FO
TITLE OF INVENTION: AMALYSIS OF GENE EXPRESSION IN BT 474
CURRENT APPLICATION NUMBER: US 0004 w0 0 8/10/203,138A
CURRENT FILING DATE: 2002-08-02
PRIOR APPLICATION NUMBER: US 60/180,312
                                                                                                                                                                                                                                           348 TAATTGTAGAATTAATTTTATAATGAAAATGCTCCAAATTTTAATGCAGGTGCTAGTAC 407
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                                                                                                                                Length 1075;
                 OTHER INFORMATION: EST_HUMAN HIT: AV739739.1, EVALUE 1.00e+00
                                                                                                                        3.3%; Score 45.8; DB 6; Length 10
41.7%; Pred. No. 0.45;
Live 0; Mismatches 402; Indels
                                                        ; OTHER INFORMATION: NT HIT: AL163201.2, EVALUE 2.00e-19 US-10-203-138A-7632
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                                                                                                                   Query Match 3.39
Best Local Similarity 41.77
Matches 287; Conservative
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APPLICANT: Molecular Dynamics, Inc.
APPLICANT: Penn, Sharron G.
APPLICANT: Penn, Sharron G.
APPLICANT: Hank, David R.
APPLICANT: Hanzel, David R.
TITLE OF INVENTION: HUWAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR TITLE OF INVENTION: HUMBER: US 4/10/203,138A
CURRENT APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2002-08-02
PRIOR PELING DATE: 03 August 2000 (36.05.00)
PRIOR FILING DATE: 03 August 2000 (36.05.00)
PRIOR APPLICATION NUMBER: US 60/207,466
PRIOR APPLICATION NUMBER: US 60/207,466
PRIOR APPLICATION NUMBER: US 60/207,466
PRIOR APPLICATION NUMBER: US 60/204,667
PRIOR PRIOR APPLICATION NUMBER: US 60/204,667
PRIOR PRIOR APPLICATION NUMBER: US 60/204,667
PRIOR PRIOR PRIOR DATE: 30 June 2000 (30.06.00)
PRIOR FILING DATE: 30 June 2000 (30.06.00)
PRIOR FILING DATE: 30 June 2000 (30.06.00)
PRIOR PRIOR PRIOR MOLECULAR Dynamics Sequence Listing Engine
PROFESSOL DATE: 307
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                                                                                                                                                                                     220 GAIGCIGGIGCTIAACCAAATCCACCIGCIACIGCTAATTTAGICACATAATGIAACGIT 279
                                                                                                                                                                                                                        280 AAATGCCCTGCTGCTACCGCAATTGCAGGTGGAGCAACAGATTATGCAGCAATAATCACA 339
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                                                                                                                 Length 1954;
                                                                                                      Score 47.4; DB 6; Length 1
Pred. No. 0.21;
0; Mismatches 171; Indels
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OTHER INFORMATION: EXPRESSED IN BT474, SIGNAL =
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Sequence 7632, Application US/10203138A
GENERAL INFORMATION:
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                                                                                                        3.48;
                                                                                                Query Match 3.4
Best Local Similarity 46.7
Matches 150; Conservative
                 ; TYPE: DNA
; ORGANISM: HUMAN
US-10-144-771-4707
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ORGANISM: HOMO
LENGTH: 1954
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      PRIOR FILING DATE: 04 February 2000 (04.02.00)
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 26 May 2000 (26.05.00)
PRIOR APPLICATION NUMBER: US 09/632,366
PRIOR PILING DATE: 03 August 2000 (03.08.00)
PRIOR FILING DATE: 03 October 2000 (03.10.00)
PRIOR FILING DATE: 20 Soctober 2000 (27.09.00)
PRIOR FILING DATE: 21 September 2000 (27.09.00)
PRIOR PRIOR FILING DATE: 21 September 2000 (27.09.00)
PRIOR APPLICATION NUMBER: US 60/236,488
PRIOR APPLICATION NUMBER: US 60/234,488
PRIOR FILING DATE: 30 June 2000 (30.06.00)
PRIOR FILING DATE: 30 June 2000 (30.06.00)
NUMBER OF SEQ ID NOS: 15438
SOCTWARE: MOLECULAR DYNAMICS SEQUENCE LISTING Engine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    y Match

Josal Similarity 41.7%; Pred. No. 0.46;

Local 287; Conservative 0; Mismatches 402;
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OTHER INFORMATION: MAP TO AL078472.1
                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 2529
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Sequence 2243, Application US/10092411A
Sequence 2243, Application US/10092411A
Sequence 2243, Application:
GENERAL INFORMATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCC
TITLE OF INVENTION: NUCLEIC ACID AND THERAPEUTICS
FILE REFERENCE: 032796-101
CURRENT APPLICATION NUMBER: US/10/092,411A
CURRENT FILING DATE: 2002-03-07
PRIOR APPLICATION NUMBER: US 09/134,001
PRIOR FILING DATE: 1998-08-13
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TITLE OF INVENTION: LANA Viral Protein Binding Sites
FILE REPERENCE: WW-07142
CURRENT APPLICATION NUMBER: US/10/194,046
CURRENT FILING DATE: 2002-07-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 45.8; DB 6;
Pred. No. 0.53;
0; Mismatches 247;
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ORGANISM: Kaposi's sarcoma-associated herpesvirus
1226 regreeregreererrerserrers 1254
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                                                                                                                                                                                                                                           ; Sequence 1, Application US/10194046; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GCTCATCCTGCTGCTCAT 2033
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US-10-092-411A-2243
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GENERAL INFORMATION:
TATLE OF INVENTION: POLYMORPHISMS IN HUMAN G-PROTEIN COUPLED
TITLE OF INVENTION: RECEPTOR GENES ASSOCIATED WITH DISEASE, METHODS OF DETECTION
TITLE OF INVENTION: AND USES THEREOF
TITLE OF INVENTION: AND USES THEREOF
CURRENT FILING DATE: 2002-12-20
PRIOR PAPLICATION NUMBER: 60/232,045
PRIOR FILING DATE: 2000-09-13
NUMBER OF SEQ ID NOS: 6961
SEQ ID NOS: 6961
SEQ ID NO 601
LENGTH: 21423
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                                                                                                                                                              Score 45.8; DB 6; Length 1
Pred. No. 0.63;
0; Mismatches 162; Indels
PRIOR APPLICATION NUMBER: US 60/064,964 PRIOR FILING DATE: 1997-11-08 PRIOR APPLICATION NUMBER: US 60/055,779 PRIOR FILING DATE: 1997-08-14 NUMBER OF SEQ ID NOS: 5676 SEQ ID NO 2243 LENGTH: 11091
                                                                                                  TYPE: DNA
; ORGANISM: Staphylococcus epidermidis
US-10-092-411A-2243
                                                                                                                                                                3.38;
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Best Local Similarity 46.99
Matches 143; Conservative
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50; Conservative
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US-09-949-004-601
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618 AGGTAAAAGTTAATGCACACCTTGTCCGGCAATTAAACCTGCTAATGTTGCTTAAGCTAC 677
                                                                                      678 TITAGGTAATGATGCTACAATAACCGCATAATGTAACGTTGCATGCCTGATGGTACTAT
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